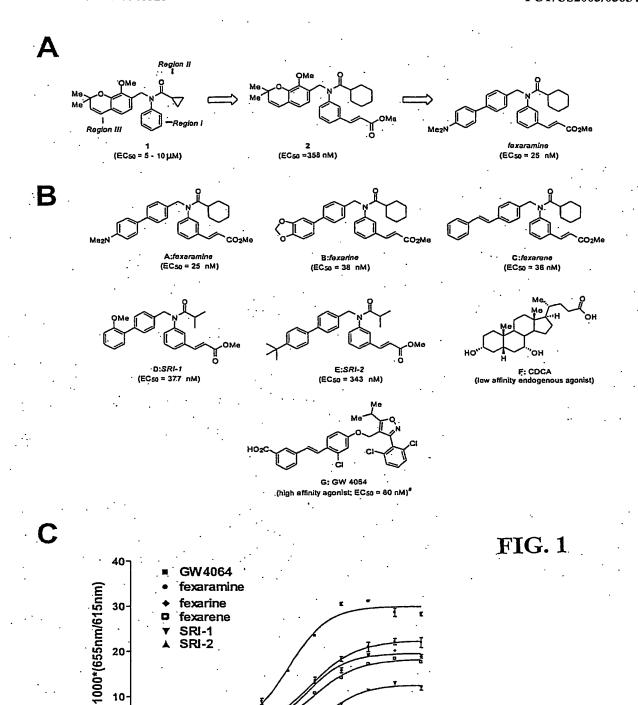
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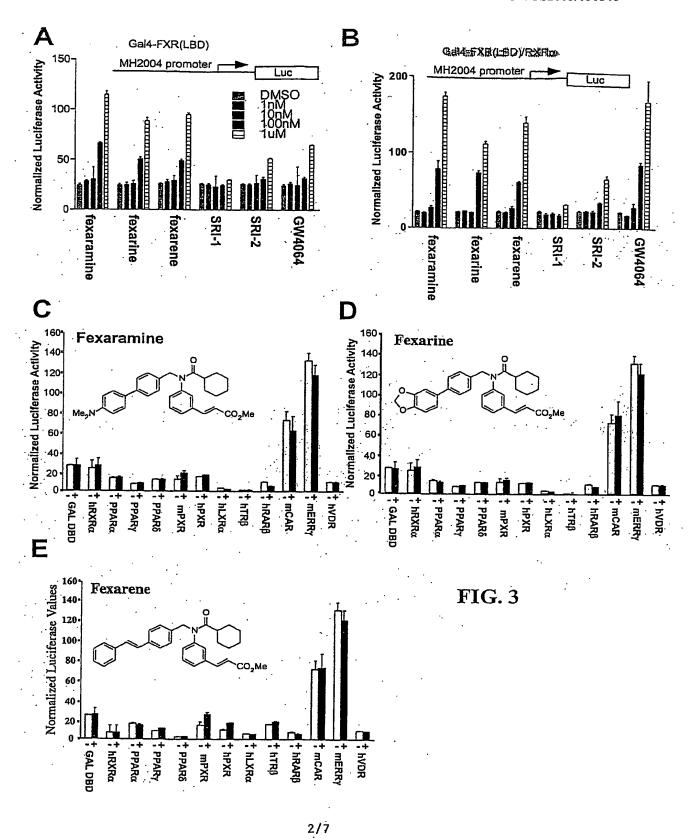
0-. -10



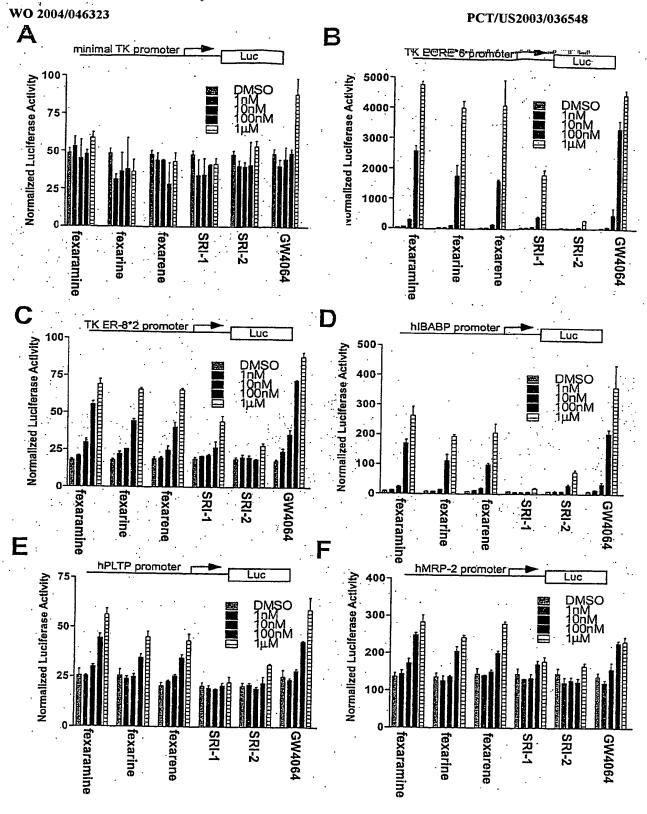
1/7

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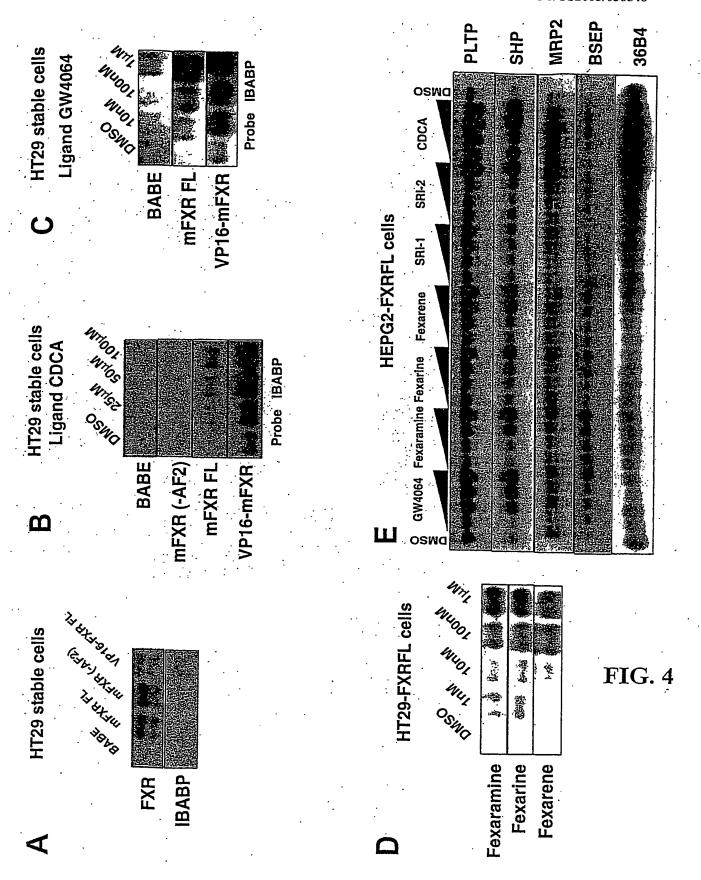
Concentration of ligands added in Log(M)



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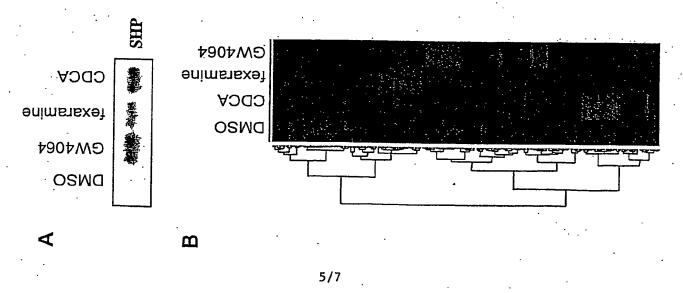
3/7



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Accession Number	Cene Description	Fold Change	104
NM_014045	apolipoprotein E	252.89	
NM_003564	transgelin 2	47.07	,
	polymerase (RNA) II (DNA directed)	45.72	
NM_001909	cathepsin D (lysosomal aspartyl protease)	42.49	-
	MGAT4B	20.06	ŀ
	WD repeat domain 18	19.03	
	Jun D proto-oncogene	16.87	,
	ribosomal protein S11	14.50	+
	FK506 binding protein 5	13.73	,
	sialyltransferase 4C	13.70	+
NM_005354	Jun D proto-oncogene	11.48	1
	glucose regulated protein, 58kD	9.03	,
	polypyrimidine tract binding protein 1	8.83	
NM 001023	ribosomai protein S20	8.58	+
NM_006579	emopamil binding protein (sterol isomerase)	7.95	+
	glucose-6-phosphatase, transport protein 1	7.85	ı
	SNARE protein	7.69	
	pre-B-cell feukemla transcription factor 2	7.49	+
NM_016504	ribosomal protein L27	7.17	+
	CED-6 protein	7.10	+
п	runt-related transcription factor 1	6.69	+
	E2F transcription factor 5, p130-binding	6.52	+
	kinase suppressor of ras	6.49	+
	FCAR (CD89):	6.45	+
	hypothetical protein FLJ12987	6.41	+
NM 001656	ADP-ribosylation factor domain protein 1	6.21	+
AK023918	novel	5.94	+
	SLC22A7	5.41	+
XP_166583	biliverdin reductase A (BLVRA)	5.30	+
AU159276	Novel	5.28	+
NM_013369	DNA (cytosine-5-)-methyltransferase 3 beta	5.23	1
NM 131836	SIM2	5.04	+
NM 007514	SLC7A2	4.64	,
NM. 002695	polymerase (RNA) II (DNA directed) gamma 2	4.85	+
AI596398	KIAA0576 protein	4.61	+
NM 004072	chemokine-like receptor 1	4.40	+
- 11	POU domain, class 4, transcription factor 1	4.00	+
NM_000369	thyrold stimulating hormone receptor	3.58	1.
	thyrold transcription factor 1	3.54	+
NM 045/42	0 = - () = - (•

FIG. 5



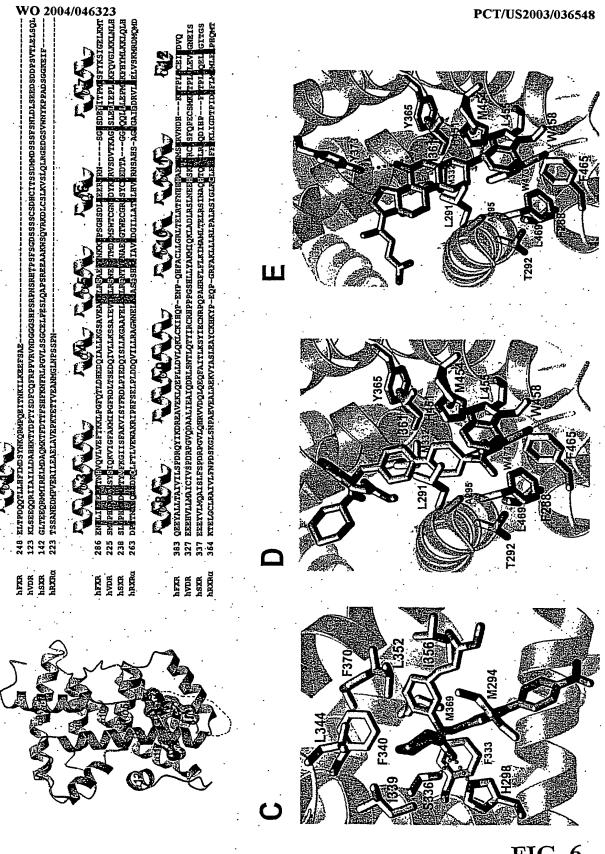


FIG. 6

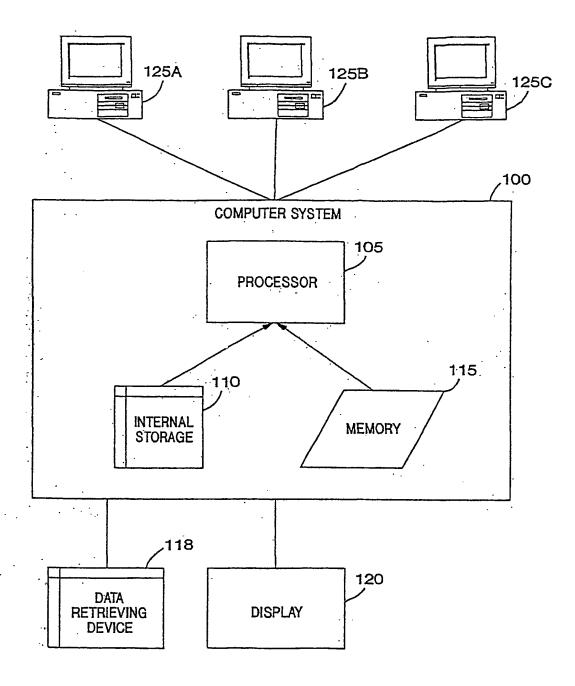


FIG. 7

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